

SEQUENCE LISTING

5 (1) GENERAL INFORMATION:

(i) APPLICANT: Arthur A. Branstrom
Donata R. Sizemore
Jerald C. Sadoff

(ii) TITLE OF INVENTION: Bacterial Delivery System

10 (iii) NUMBER OF SEQUENCES: 8

(iv) CORRESPONDENCE ADDRESS:

15 (A) ADDRESSEE: John Moran
(B) STREET: USA MPMC - MCMR-JA
(C) CITY: FORT DETRICK, FREDERICK
(D) STATE: MARYLAND
(E) COUNTRY: USA
(F) ZIP: 21702-5012

20 (v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: Apple Macintosh
(C) OPERATING SYSTEM: Macintosh 7.0
(D) SOFTWARE: Microsoft Word

25 (vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:

30 (vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE:

35 (viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Moran, John
(B) REGISTRATION NUMBER: 26,313
(C) REFERENCE/DOCKET NUMBER:

40 (ix) TELECOMMUNICATION INFORMATION

(A) TELEPHONE: (301) 619-2065
(B) TELEFAX: (301) 619-7714

45 (2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

50 (A) LENGTH: 1674 base pairs
(B) TYPE: Nucleic acid
(C) STRANDEDNESS: Double
(D) TOPOLOGY: Linear

SEQUENCE DESCRIPTION: SEQ ID NO: 1

	TCCATAATCA GGATCAATAA AACTGCTGCA GAAATGATTT	40
	CATTCATAAC TCAAATTCCC TGATAATTGC CGCGGACTTT	80
5	CTGCGTGCTA ACAAAGCAGG ATAAGTCGCA TTAATCATGG	120
	CTTCGCTATC ATTGATTAAT TTCACTTGCG ACTTTGGCTG	160
	CTTTTTGTAT GGTGAAAGAT GTGCCAAGAG GAGACCGGCA	200
	CATTTATACA GCACACATCT TTGCAGGAAA AAAACGCTTA	240
	TGAAAAATGT TGGTTTTATC GGCTGGCGCG GTATGGTCGG	280
10	CTCCGTTCTC ATGCAACGCA TGGTTGAAGA GCGCGACTTC	320
	GACGCCATTC GCCCTGTCTT CTTTCTACT TCTCAGCTTG	360
	GCCAGGCTGC GCCGTCTTTT GGCGGAACCA CTGGCACACT	400
	TCAGGATGCC TTTGATCTGG AGGCGCTAAA GGCCCTCGAT	440
	ATCATTGTGA CCTGTCAGGG CGGCGATTAT ACCAACGAAA	480
15	TCTATCCAAA GCTTCGTGAA AGCGGATGGC AAGGTTACTG	520
	GATTGACGCA GCATCGTCTC TGCGCATGAA AGATGACGCC	560
	ATCATCATTG TTGACCCCGT CAATCAGGAC GTCATTACCG	600
	ACGGATTAAA TAATGGCATC AGGACTTTTG TTGGCGGTAA	640
	CTGTACCGTA AGCCTGATGT TGATGTCGTT GGGTGGTTTA	680
20	TTGCGCAATG ATCTTGTTGA TTGGGTGTCC GTTGCAACCT	720
	ACCAGGCCGC TTCCGGCGGT GGTGCGCGAC ATATGCGTGA	760
	GTTATTAACC CAGATGGGCC ATCTGTATGG CCATGTGGCA	800
	GATGAACTCG CGACCCCGTC CTCTGCTATT CTCGATATCG	840
	AACGCAAAGT CACAACCTTA ACCCGTAGCG GTGAGCTGCC	880
25	GGTGGATAAC TTTGGCGTGC CGCTGGCGGG TAGCCTGATT	920
	CCGTGGATCG ACAAACAGCT CGATAACGGT CAGAGCCGCG	960
	AAGAGTGGA AGGGCAGGCG GAAACCAACA AGATCCTCAA	1000

	CACATCTTCC GTAATTCCGG TAGATGGTTT ATGTGTGCGT	1040
	GTCGGGGCAT TGCCTGCCA CAGCCAGGCA TTCCTATTA	1080
	AATTGAAAAA AGATGTGTCT ATTCCGACCG TGGAAGAACT	1120
	GCTGGCTGCG CACAATCCGT GGGCGAAAGT CGTTCCGAAC	1160
5	GATCGGGAAA TCACTATGCG TGAGCTAACC CCAGCTGCCG	1200
	TTACCGGCAC GCTGACCACG CCGGTAGGCC GCCTGCGTAA	1240
	GCTGAATATG GGACCAGAGT TCCTGTCAGC CTTTACCGTG	1280
	GGCGACCAGC TGCTGTGGGG GGCCGCGGAG CCGCTGCGTC	1320
	GGATGCTTCG TCAACTGGCG TAATCTTTAT TCATTAAATC	1360
10	TGGGGCGCGA TGCCGCCCCT GTTAGTGCGT AATACAGGAG	1400
	TAAGCGCAGA TGTTTCATGA TTTACCGGGA GTTAAATAGA	1440
	GCATTGGCTA TTCTTTAAGG GTGGCTGAAT ACATGAGTAT	1480
	TCACAGCCTT ACCTGAAGTG AGGACGACGC AGAGAGGATG	1520
	CACAGAGTGC TGCGCCGTTT AGGTCAAAAA AATGTCACAA	1560
15	CCAGAAGTCA AAAATCCAAT TGGATGGGGT GACACAATAA	1600
	AACAGGAAGA CAAGCATGTC CGATCGTATC GATAGAGACG	1640
	TGATTAACGC GCTAATTGCA GGCCATTTTG CGGA	1674

(3) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1121 base pairs
- (B) TYPE: Nucleic acid
- (C) STRANDEDNESS: Double
- (D) TOPOLOGY: Linear

(ii) MOLECULE TYPE: Other nucleic acid

(A) DESCRIPTION: The *E. coli asd* gene coding for β -aspartic semialdehyde dehydrogenase identified in SEQ ID NO:1 was modified by deleting 553 bp from position 439 to 991.

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2

TCCATAATCA GGATCAATAA AACTGCTGCA GAAATGATTT

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(4) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: Nucleic acid
- (C) STRANDEDNESS: Double
- (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3
AGATCTCCCT GATAATTGCC GC

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(5) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 26 base pairs
- (B) TYPE: Nucleic acid
- (C) STRANDEDNESS: Double
- (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4
AGATCTCGCT TACTCCTGTA TTACGC

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(6) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: Nucleic acid
- (C) STRANDEDNESS: Double
- (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5
CGAGGGCCTT TAGCGCCTCC

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(7) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: Nucleic acid
- (C) STRANDEDNESS: Double
- (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6

GATCCTCAAC ACATCTTCCG

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5 (8) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 22 base pairs
(B) TYPE: Nucleic acid
(C) STRANDEDNESS: Double
(D) TOPOLOGY: Linear

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7

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GAGCTCCCCT GATAATTGCC GC

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(9) INFORMATION FOR SEQ ID NO:8:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 26 base pairs
(B) TYPE: Nucleic acid
(C) STRANDEDNESS: Double
(D) TOPOLOGY: Linear

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8

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GTCGACCGCT TACTCCTGTA TTACGC

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